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Formation of a σ -alkane Complex and a Molecular Rearrangement in the Solid-State: $[\text{Rh}(\text{Cyp}_2\text{PCH}_2\text{CH}_2\text{PCyp}_2)(\eta^2:\eta^2\text{-C}_7\text{H}_{12})][\text{BAr}^{\text{F}}_4]$

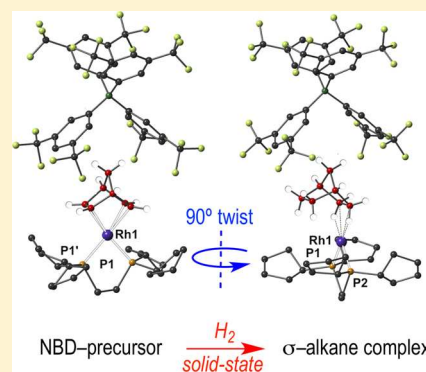
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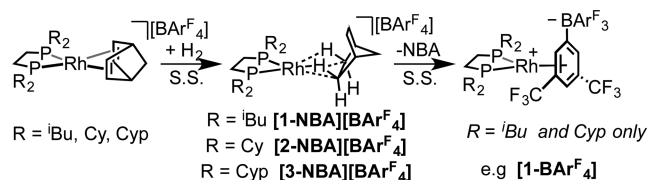
S Supporting Information

ABSTRACT: Addition of H_2 to the precursor $[\text{Rh}(\text{Cyp}_2\text{PCH}_2\text{CH}_2\text{PCyp}_2)(\eta^2:\eta^2\text{-C}_7\text{H}_8)][\text{BAr}^{\text{F}}_4]$ gives the σ -alkane complex $[\text{Rh}(\text{Cyp}_2\text{PCH}_2\text{CH}_2\text{PCyp}_2)(\eta^2:\eta^2\text{-C}_7\text{H}_{12})][\text{BAr}^{\text{F}}_4]$ by a single-crystal to single-crystal reaction, as characterized by X-ray crystallography, SSNMR spectroscopy, and periodic DFT. An unexpected rearrangement of the $\{\text{Rh}(\text{L}_2)\}^+$ fragment is revealed.



Complexes in which an alkane acts as a ligand to a metal center using its C–H bonding electrons, so-called σ -alkane complexes, are of considerable interest due to their role as intermediates in C–H activation processes and the challenges involved with their synthesis and characterization.¹ As C–H bonds are strong and non-nucleophilic, alkanes are very poor ligands, and direct observation of σ -alkane complexes has generally been limited to very low temperature in situ solution spectroscopic techniques.² We have recently reported that they may be prepared by solid/gas single-crystal to single-crystal transformations (Scheme 1),³ enabling structural

Scheme 1. Synthesis of σ -Alkane Complexes^a



^aS. S. = solid-state.

characterization by single crystal X-ray crystallography. Addition of H_2 to the diene precursor $[\text{Rh}(\text{R}_2\text{PCH}_2\text{CH}_2\text{PR}_2)(\eta^2:\eta^2\text{-NBD})][\text{BAr}^{\text{F}}_4]$ (NBD = norbornadiene, Ar^F = 3,5-(CF₃)₂C₆H₃) leads to the generation of $[\text{Rh}(\text{R}_2\text{PCH}_2\text{CH}_2\text{PR}_2)(\eta^2:\eta^2\text{-NBA})][\text{BAr}^{\text{F}}_4]$ (NBA = norbornane; R = ^tBu, [1-NBA][BAr^F₄]; R = Cy, [2-NBA][BAr^F₄]). When R = ^tBu, rapid loss of NBA in the solid-state at 298 K gives the [BAr^F₄]-coordinated zwitterion [1-BAr^F₄].⁴ In contrast, when R = Cy,

the σ -alkane complex is stable in the solid state for months at 298 K.⁵ The precise factors that are responsible for these differences are currently not fully resolved. Both species feature encapsulation of the cation by an octahedral arrangement of [BAr^F₄][−] anions, which provides a well-defined lattice environment, while the buttressing Cy groups in [2-NBA][BAr^F₄] may install a kinetic penalty toward rearrangement in the solid state.^{4,5} We now report the synthesis of $[\text{Rh}(\text{Cyp}_2\text{PCH}_2\text{CH}_2\text{PCyp}_2)(\eta^2:\eta^2\text{-NBA})][\text{BAr}^{\text{F}}_4]$ using a cyclopentyl (Cyp)-substituted chelating phosphine⁶ which was designed to sit between the steric profiles of flexible ^tBu and rigid/tall Cy. This leads to not only a relatively stable σ -alkane complex but also an unexpected metal-fragment reorientation within the [BAr^F₄][−] anion cavity while retaining the integrity of the single crystal.

The precursor complex $[\text{Rh}(\text{Cyp}_2\text{PCH}_2\text{CH}_2\text{PCyp}_2)(\eta^2:\eta^2\text{-NBD})][\text{BAr}^{\text{F}}_4]$ ([3-NBD][BAr^F₄]) was prepared as for [2-NBD][BAr^F₄].⁵ A single-crystal X-ray diffraction study at 150 K⁷ (Figure 1A) demonstrated crystallographically imposed C₂ rotational symmetry with Rh1 and NBD bridge methylene (C7) occupying special positions. The Cyp groups are disordered, while the long-range structure was found to be modulated (Supporting Information),⁸ which was modeled with a *q* vector of (0,0.3433,0).⁹ The [BAr^F₄][−] anions are organized so that each cation is contained within a pseudo-octahedral cage of anions, and the overall motif and structural

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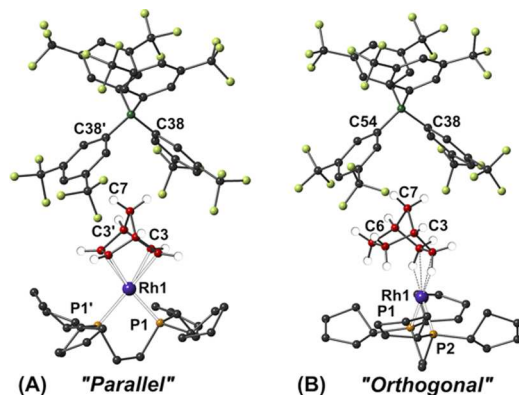


Figure 1. Solid-state structures (ball and stick) relative to a local $[\text{BARF}_4]^-$ anion: (A) $[\text{3-NBA}][\text{BARF}_4]$, average structure; (B) $[\text{3-NBD}][\text{BARF}_4]$. Selected interplane angles: $\text{C38}', \text{C38}, \text{Rh1}/\text{Rh1}, \text{P1}, \text{P1}' = 6.39(17)^\circ$; $\text{C38}, \text{C54}, \text{Rh1}/\text{Rh1}, \text{P1}, \text{P2} = 88.12(12)^\circ$.

metrics are very similar to those reported for $[\text{2-NBD}][\text{BARF}_4]$. In particular the NBD ligand sits in a cleft between two $[\text{BARF}_4]^-$ aryl groups so that the C7-methylene protons point toward the centers of two of the aromatic rings. In the 298 K $^{31}\text{P}\{\text{H}\}$ solid-state NMR (SSNMR) spectrum two distinct, but very close, phosphine environments are observed (δ 66.2, J_{RhP} 149 Hz; δ 65.7, J_{RhP} 151 Hz), consistent with PCyp₂ disorder and/or the modulated structure. In the $^{13}\text{C}\{\text{H}\}$ SSNMR spectrum four signals observed between δ 90 and 55 are assigned to the NBD fragment under C_2 symmetry. The disorder/modulation was not resolved in this spectrum.

Addition of H_2 to a single-crystal sample of $[\text{3-NBD}][\text{BARF}_4]$ (2 bar, 10 min, 298 K) led to the quantitative formation of the σ -alkane complex $[\text{Rh}(\text{Cyp}_2\text{PCH}_2\text{CH}_2\text{PCyp}_2)(\eta^2\text{-}\eta^2\text{-NBA})][\text{BARF}_4]$ ($[\text{3-NBA}][\text{BARF}_4]$) (Scheme 1). Crystallinity is retained in this process, and the resulting structure determined by single-crystal X-ray diffraction (100 K, $R(2\sigma) = 7.0\%$) clearly shows a saturated NBA fragment interacting with the Rh center through two 3c-2e $\text{Rh}\cdots\text{H}-\text{C}$ interactions, in which the relevant hydrogen atoms were located but refined using a riding model (Figure 2). Although there are no significant changes to the arrangement of the $[\text{BARF}_4]^-$ anions, this

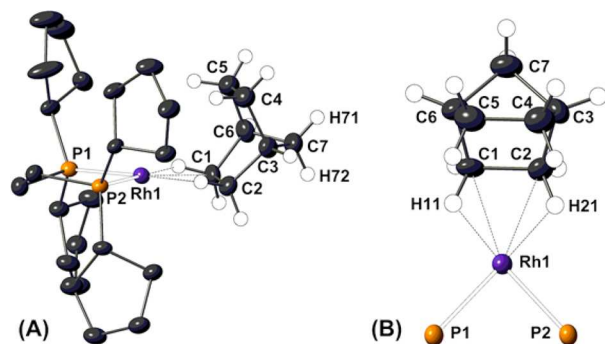


Figure 2. (A) Solid-state structure of $[\text{3-NBA}][\text{BARF}_4]$, with the cation and selected H atoms shown (100 K, 50% displacement ellipsoids). (B) Alternate view of $[\text{3-NBA}][\text{BARF}_4]$, with P-alkyl groups removed. Selected bond lengths (\AA): $\text{Rh1}-\text{C1}$ 2.388(5), $\text{Rh1}-\text{C2}$ 2.392(5), $\text{Rh1}-\text{P1}$ 2.207(1), $\text{Rh1}-\text{P2}$ 2.206(1), $\text{C1}-\text{C2}$ 1.550(7), $\text{C4}-\text{C5}$ 1.541(9). Computed bond lengths (\AA ; periodic DFT, PBE-D3): $\text{Rh1}-\text{C1}$ 2.425, $\text{Rh1}-\text{C2}$ 2.420, $\text{Rh1}-\text{P1}$ 2.219, $\text{Rh1}-\text{P2}$ 2.215, $\text{C1}-\text{C2}$ 1.561, $\text{C4}-\text{C5}$ 1.565, $\text{Rh1}\cdots\text{H11}$ 1.895, $\text{Rh1}\cdots\text{H21}$ 1.886, $\text{C1}-\text{H11}$ 1.154, $\text{C2}-\text{H21}$ 1.154.

transformation proceeds with loss of the crystallographically imposed C_2 rotational symmetry: $[\text{3-NBD}][\text{BARF}_4]$, C_2/c , $Z = 4$; $[\text{3-NBA}][\text{BARF}_4]$, $P2_1/c$, $Z = 4$.¹⁰ Space group changes have been noted in other single-crystal transformations.^{5,11} The alkane ligand binds to the $\{\text{Rh}(\text{L}_2)\}^+$ fragment through two C–H σ -interactions, resulting in a Rh(I) square-planar, d^8 metal center coordination motif. The Rh–C1/C2 distances (2.388(5) and 2.392(5) \AA , respectively) are the same, within error, as for $[\text{2-NBA}][\text{BARF}_4]$ (2.389(3) and 2.400(3) \AA). The C1–C2 and C4–C5 distances (1.550(7) and 1.541(9) \AA) signal the hydrogenation of the diene to form C–C bonds. The Rh–P distances (2.207(1) and 2.206(1) \AA) are shorter than those in $[\text{3-NBD}][\text{BARF}_4]$ (2.2712(12) \AA), reflecting the weaker trans influence C–H σ interactions. The Cyp groups showed some disorder.

The changes upon hydrogenation were also probed by the optimization of the extended solid-state structure with periodic DFT calculations at the PBE-D3 level. These provided excellent agreement both for the molecular geometries of the Rh cations (see the caption of Figure 2 for $[\text{3-NBA}][\text{BARF}_4]$ and the Supporting Information for $[\text{3-NBD}][\text{BARF}_4]$) and for the extended structure (see Figure S15 in the Supporting Information). Moreover, the calculations provide further insight into the $\text{Rh}\cdots\text{alkane}$ interaction, with short average computed $\text{Rh}\cdots\text{H11}/\text{H21}$ distances (1.89 \AA) and elongated C1–H11 and C2–H21 distances (1.15 \AA). In contrast, the C1–H12 and C2–H22 bonds exhibit standard distances of 1.10 \AA , consistent with an absence of any interaction with Rh in that case. The $\text{Rh}\cdots\text{H}-\text{C}$ σ -interaction is confirmed by a quantum theory of atoms in molecules (QTAIM) study that identifies $\text{Rh}\cdots\text{H11}$ and $\text{Rh}\cdots\text{H21}$ bond paths with reduced electron densities for the C1–H11 and C2–H21 bond critical points in comparison to the C1–H12 and C2–H22 bonds.⁹

Consideration of the relationship between the cation and the proximal capping $[\text{BARF}_4]^-$ anion (Figure 1) shows that the NBA ligand in $[\text{3-NBA}][\text{BARF}_4]$ adopts an orientation very similar to that of the NBD ligand in $[\text{3-NBD}][\text{BARF}_4]$, in particular the orientation of the C7-methylene protons that are still directed toward the aryl rings. In contrast the $\{\text{Rh}(\text{L}_2)\}^+$ fragment has undergone a ca. 90° twist with respect to the precursor as defined by, for example, the interplane angle $\text{C38}, \text{C54}, \text{Rh1}/\text{Rh1}, \text{P1}, \text{P2} = 88.12(12)^\circ$ (Figure 1). We have previously noted¹² that the octahedral cavity described by the anions accommodates a variety of structural changes associated at the metal cation. Clearly it can also accommodate significant movement of the $\{\text{Rh}(\text{L}_2)\}^+$ fragment. Whether this occurs in concert with hydrogenation, or immediately afterward, is opaque to experiment. This situation is different from that observed in the transformations that afforded $[\text{2-NBA}][\text{BARF}_4]$, in which it is the organic fragment that responds by moving rather than $\{\text{Rh}(\text{L}_2)\}^+$.⁵ Isomerization, or dynamic processes, in single-crystal organometallics have been reported.¹³

Computationally the alternative form of $[\text{3-NBA}][\text{BARF}_4]$ was probed by rotating one complete cation within the unit cell and reoptimizing the structure (Scheme 2). This gives a local “parallel” structure analogous to that seen within $[\text{2-NBA}][\text{BARF}_4]$ and equivalent to the structure that would be formed if, upon hydrogenation, the NBD ligand in $[\text{3-NBD}][\text{BARF}_4]$ underwent rotation rather than the $\{\text{Rh}(\text{L}_2)\}^+$ cation. For $[\text{3-NBA}][\text{BARF}_4]$ the parallel structure is ca. 14 kcal/mol above the optimized “orthogonal” form. The parallel structure is therefore not viable thermodynamically, but it may be kinetically accessible, 14 kcal/mol representing a lower limit to the

Notes

The authors declare no competing financial interest.

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- (10) Initial data collection at 150 K revealed a poorer structure with significantly larger displacement ellipsoids associated with the NBA. There is only a 0.9% change in unit cell volume between [3-NBD][BAr^F₄] and [3-NBA][BAr^F₄] at 150 K: $V = 6142.10(16)$, $6197.3(3)$ Å³, respectively.
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